10,536 LCMS hours (charged: 5,912 hours) not including Quality Control and most implementation/tests.

3,169 digestions (charged: 1,731).

413 Proteomics projects (240 internal) and 179 Small Molecule projects (153 internal)

179 individual users (Internal 119/External 60) from 79 different labs (Internal 44/External 35). Walkup instrument usage is not included.

FY21 subsidy: 34.4%.
Projected FY22 subsidy: 30% (based on first 5 months).

29% of all mass spec hours were used by the Metabolomics Platform.

Median and average turn-around time: 9 and 15 days

10 letters of support for grants and applications.

12 co-authored publications and 7 acknowledgements.